

AMENDMENTS TO THE SPECIFICATION

Please amend the paragraph beginning on page 35, line 28, as follows:

Bioinformatics analyses. Bioinformatics analysis was performed with the NCBI Human Genome databases (available on the Web at ncbi.nlm.nih.gov/genome/guide/human) and the Celera Discovery System (available online at cds.celera.com/cds~~http://eds.celera.com/eds~~). Gene finding was performed with BLASTX (available on the Web at ncbi.nlm.nih.gov/BLAST) and GENESCAN and FGENES software (GeneMachine, DIR, NIH, [available online at genome.nhgri.nih.gov/genemachine/](http://genome.nhgri.nih.gov/genemachine/)~~http://genome.nhgri.nih.gov/genemachine/~~). SNPs were developed using the SNP database (~~available on the Web at ncbi.nlm.nih.gov/SNP/~~)[\(available on the Web at ncbi.nlm.nih.gov/SNP/\)](http://ncbi.nlm.nih.gov/SNP/).